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DPP4 Inhibition Ameliorates Cardiac Function by Blocking the Cleavage of HMGB1 in Diabetic Mice After Myocardial Infarction

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Summary

High mobility group box 1 (HMGB1), a ubiquitous DNA-binding protein, promotes angiogenesis and tissue repair, resulting in restored cardiac function after myocardial infarction (MI). Although dipeptidyl peptidase 4 (DPP4) degrades certain peptides, it remains unclear as to whether HMGB1 is a substrate of DPP4 and whether DPP4 inhibition prevents the cleavage of HMGB1.

In transgenic mice with cardiac-specific overexpression of HMGB1 (TG) and wild-type mice (WT), a diabetic state was induced by streptozotocin, and MI was created by ligation of the left anterior descending coronary artery. To inhibit DPP4 activity, a DPP4 inhibitor anagliptin was used. The plasma levels of HMGB1, infarct size, echocardiographic data, angiogenesis, and vascular endothelial growth factor (VEGF) expression in the peri-infarct area were compared among non-diabetic MI WT/TG, diabetic MI WT/TG, and anagliptin-treated diabetic MI WT/TG mice.

DPP4 activity was increased in the diabetic state and blocked by anagliptin administration. The HMGB1 plasma levels were reduced in the diabetic TG compared with the non-diabetic TG mice, but DPP4 inhibition with anagliptin increased HMGB1 plasma levels in the diabetic TG mice. The infarct area was significantly larger in the diabetic TG than in the non-diabetic TG mice, and it was reduced by DPP4 inhibition. Cardiac function, angiogenesis, and VEGF expression were impaired in the diabetic TG mice, but they were ameliorated by the DPP4 inhibition to levels similar to those found in the non-diabetic TG mice.

The DPP4 inhibitor ameliorated cardiac function by inhibiting the inactivation of HMGB1 in diabetic mice after MI.

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Key words: Angiogenesis, Ischemic heart disease, Diabetes mellitus

Acute myocardial infarction (MI) is an emergent disease caused by occlusion of the coronary artery. The morbidity and mortality of MI remain high despite the advancement of emergency medical services and the development of optimal therapies, including pharmacological treatment and percutaneous coronary intervention, and surgical technique.¹⁾ Infarct size is known as an important prognostic predictor; thus, many studies of ischemic heart disease have been performed to determine how to limit the extent of infarction, focusing on the mechanisms of collateral development, ischemic reperfusion injury and ischemic preconditioning, among others.²⁻⁴⁾

High mobility group box 1 (HMGB1) is a ubiquitous non-histone DNA-binding protein that mediates gene transcription.⁵⁻⁷⁾ HMGB1 secreted from necrotic cells into the extracellular space triggers tissue repair as a cytokine.⁸⁻¹¹⁾ Moreover, several studies have proven that HMGB1 is involved in limiting the size of the infarct area and preserv-

ing cardiac function by promoting angiogenesis.^{12,13)} In addition, we have previously demonstrated that HMGB1 promotes angiogenesis and tissue repair by enhancing mobilization and differentiation of endothelial progenitor cells from bone marrow, resulting in preserving cardiac function after MI.¹⁴⁾ These data suggest that HMGB1 is a possible cardio-protective protein.

Diabetes mellitus (DM) is a major coronary risk factor, and MI patients with DM are likely to have a poor prognosis compared with those without DM.¹⁵⁾ Over the past few years, many researchers have shown an interest in the pleiotropic effect of the inhibition of dipeptidyl peptidase 4 (DPP4), whose activity is upregulated in a diabetic state.^{16,17)} A DPP4 inhibitor has an anti-diabetic effect by suppressing the degradation of incretin hormones such as glucagon-like peptide-1 (GLP-1) and gastrointestinal peptide (GIP).¹⁸⁾ It has recently been reported that DPP4 degrades not only incretin hormones but also some

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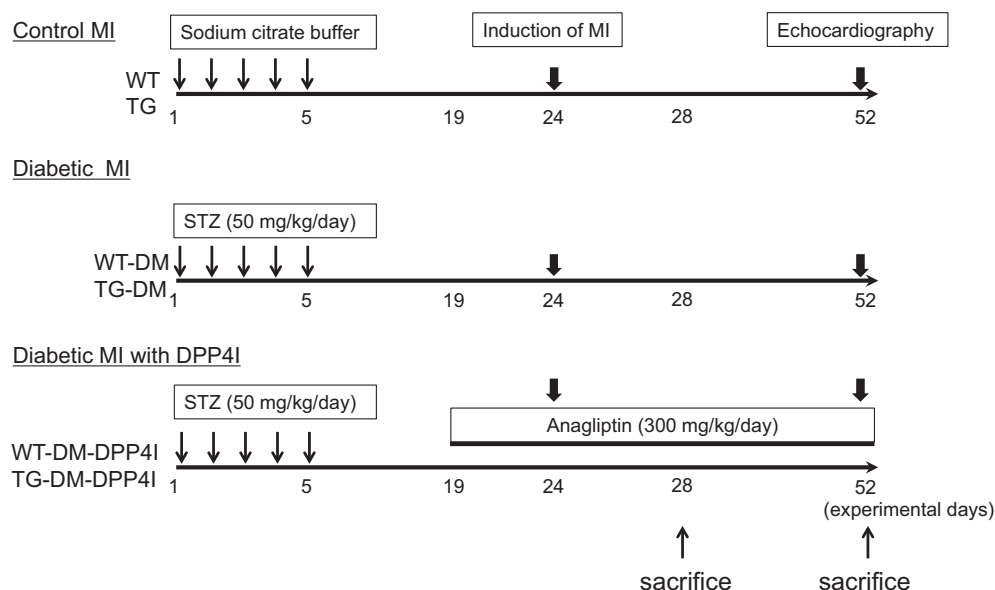


Figure 1. The schemas of the experimental protocols for each of the six groups used in this study. The mice were divided into six groups (WT, WT-DM, WT-DM-DPP4I, TG, TG-DM, and TG-DM-DPP4I) and experimental myocardial infarction (MI) was induced in all groups. Anagliptin administration was started at five days before induction of experimental MI and then continued through the experimental periods in the WT-DM-DPP4I and TG-DM-DPP4I groups.

peptides with a role in the regulation of the cardiovascular system, such as stromal-cell-derived factor-1 (SDF-1), B-type natriuretic peptide (BNP) and substance P, as well as other peptides.¹⁹⁾ Although some reports have claimed that HMGB1 can also be degraded by DPP4,²⁰⁾ it still remains unclear as to whether DPP4 cleaves HMGB1 *in vivo*.

In the present study, we tested our hypothesis that the inhibition of DPP4 restores HMGB-1-induced angiogenesis and tissue repair in diabetic mice after MI.

Methods

Animals and ethical statements: Transgenic male mice with cardiac-specific HMGB1 overexpression (TG mice) on a BDF-1 background¹²⁾ and wild-type littermate male mice (WT mice) (10-12 weeks of age) were used for the experiments. All mice genotypes were confirmed by PCR analyses of tail DNA before the experiments. The mice were housed under pathogen-free conditions in isolated cages on a 12-hour light/dark cycle. They could freely access standard rodent food and water.

The investigations conformed to the Guide for the Care and Use of Laboratory Animals, published by the US National Institutes of Health (NIH publication, 8th Edition, 2011). The Fukushima Medical Research Committee approved our research protocol. All animal experiments were performed in accordance with the guidelines of the Fukushima Medical University Animal Research Committee, with efforts made to minimize the suffering of the animals.

Experimental protocol: The experimental protocol is shown in Figure 1. We divided the mice into six groups; control MI groups (WT and TG), diabetic MI groups

(WT-DM and TG-DM), and diabetic MI treated with DPP4 inhibitor groups (WT-DM-DPP4I and TG-DM-DPP4I). To induce a diabetic state, streptozotocin (STZ, 50 mg/kg/day, Sigma, St. Louis, MO, USA), which was dissolved in 0.1 M sodium citrate buffer (pH 4.5) as described in previous reports,²¹⁾ was administered for five consecutive days by intraperitoneal injection. Sodium citrate buffer without STZ was administered to control MI group for five consecutive days by intraperitoneal injection. Two weeks after injection of STZ, whole blood glucose levels were measured by a glucometer (GLUCOCARD™, ARKRAY Inc., Kyoto, Japan). A diabetic state was defined as when blood glucose levels were elevated to more than 16 mmol/L (288 mg/dL).²¹⁾ Experimental MI was induced as previously described^{12,14)} in all groups at day 24. Briefly, the mice under general anesthesia by intraperitoneal injection of tribromoethanol (0.25 mg/g of body weight) were intubated with a 20-gauge polyethylene catheter and ventilated using a rodent ventilator (Shinano Manufacturing, Tokyo, Japan). A chest incision was performed at the level of the fourth rib and along the left sternal border, and the left anterior descending coronary artery (LAD) was ligated with 8-0 prolene sutures. Induction of MI was considered successful when the color of the anterior wall, including the apical portion, turned pale. After LAD ligation, the chest wall was closed using 6-0 nylon sutures. Western blotting was performed on the six groups and two sham groups (WT-sham and TG-sham). The sham operation procedure was the same as induction of MI except for LAD ligation. In the WT-DM-DPP4I and TG-DM-DPP4I groups, administration of a DPP4 inhibitor, anagliptin (supplied by SANWA KAGAKU KENKYUSHO CO. LTD, Aichi, Japan), via the oral route was

started at five days before MI induction, and was then continued throughout the experimental periods. Anagliptin (300 mg/kg/day) was dissolved in drinking water. The dose of anagliptin was determined from previous reports.²²⁻²⁴⁾ In order to determine whether the dose of anagliptin (300 mg/kg/day) was adequate to inhibit DPP4 activity, DPP4 activity was measured as described in the next section, and we found DPP4 activity in this study was inhibited in nearly equal level compared with those of previous experiments (80-90% inhibition of DPP4 activity).^{22,25,26)} Echocardiography was performed in order to evaluate the cardiac function of the mice at day 52 (four weeks after MI induction). To obtain mice heart tissue samples, sacrifice was performed at one week and four weeks after induction of MI in each group.

Fluorometric assay for measurement of DPP4 activity: Plasma DPP4 activities at baseline, two weeks after STZ injection, five days after starting administration of anagliptin, and four weeks after MI operation, were measured in whole blood using a Fluorometric assay kit based on fluorescent substrates (BioVision Inc, Milpitas, CA, USA) according to the lab manual.

Enzyme-linked immunosorbent assay (ELISA): Plasma levels of HMGB1 at four days after induction of MI in each group were measured using an ELISA kit (Shino Test Corporation, Tokyo, Japan), according to the lab manual.^{12,14)}

Echocardiographic measurements: Transthoracic echocardiography was performed under light inhaled anesthesia with isoflurane in each group at four weeks after induction of MI using a Vevo 2100 High-Resolution In Vivo Imaging System (Visual Sonics Inc., Toronto, Canada) with a high-resolution 40-MHz imaging transducer.²⁷⁾ With the use of the M-mode images, interventricular septal thickness, posterior wall thickness, left ventricular end-diastolic dimension (LVDd), and left ventricular end-systolic dimension (LVDs) were measured. The percentage of left ventricular fractional shortening (FS) was calculated as $100 \times ((LVDd - LVDs) / LVDd)$.

Histological examinations: To observe morphological changes of heart tissue, mice from all groups were sacrificed by cervical dislocation at four weeks after induction of MI. The heart was excised and weighed after flushing with $1 \times$ PBS buffer. The paraffin-embedded heart tissues of each group were prepared and sliced serially at the papillary muscle level. The obtained slices were then stained with hematoxylin-eosin and Masson's trichrome stains. The infarct size was expressed as the ratio of the infarct length to the total left ventricular length calculated by averaging the endo- and epi-cardial perimeters.^{12,14)} The above analyses of the infarct size were performed using Image J software (NIH, Bethesda, MD, USA).

On immunohistochemical analysis, the paraffin sections of the heart tissue at four weeks after MI were stained by anti-platelet endothelial cell adhesion molecule (PECAM-1) antibodies (Santa Cruz, CA, USA) to identify endothelial cells and by anti- α smooth muscle actin (α SMA) antibodies (Santa Cruz, CA, USA) to identify α SMA positive cells. The paraffin sections of the heart tissue were stained with horseradish peroxidase-conjugated secondary antibodies (Histofine Simple Stain

Mouse MAX PO, Nichirei Bioscience Inc., Tokyo, Japan) and diaminobenzidine tetrahydrochloride, then counterstained with hematoxylin. To assess the number of PECAM-1- and α SMA-positive cells as indicators of angiogenesis, digital photomicrographs of the border zone (1-2 mm from the edge of the infarct area) were obtained, and the numbers of PECAM-1- and α SMA-positive cells in the border zone were counted in a randomly selected high-power field (HPF, $\times 400$) by two independent researchers. The counts were repeated ten times, and the number of PECAM-1- and α SMA-positive cells were obtained by calculating the average of the ten measurements.^{12,14)}

Western blotting: The heart tissue samples were obtained from each group at four days after induction of MI. The total protein was extracted from the snap-frozen left ventricle using Cell Lysis Buffer (Cell Signaling Technology, Beverly, MA) with Protease Inhibitor Cocktail (BD Biosciences, San Jose, CA, USA).²⁷⁾ The protein concentration of the myocardial sample was determined by protein assay (DC protein assay kit, Bio-Rad Laboratories, Hercules, CA). Equal amounts (20 μ g) of the protein samples were subjected to electrophoresis onto sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE, 10%) and transferred onto polyvinylidene difluoride membranes (ATTO Co., Tokyo, Japan). The signals from the immunoreactive bands were visualized by an Amersham ECL system (Amersham Pharmacia Biotech UK Ltd., Buckinghamshire, UK) and quantified using Image J Software (NIH). The antibodies used in the present study were rabbit polyclonal anti-vascular endothelial growth factor (VEGF) antibody (used at dilution of 1:1000, Santa Cruz) and mouse polyclonal anti- β actin antibody (1:5000, Santa Cruz). VEGF expression was normalized by β actin.

Statistical analysis: All data were presented as mean \pm standard error. Statistical significance was evaluated using one-way analysis of variance for comparisons among the six groups, followed by multiple comparisons with the Bonferroni post hoc test. A $P < 0.05$ was considered statistically significant. All analyses were performed using a statistical software package (SPSS ver. 22.0, IBM, Armonk, NY, USA).

Results

Measurements of DPP4 enzyme activity: We first confirmed whether the DPP4 activity was suppressed by oral administration of anagliptin (300 mg/kg/day). Whole blood samples were obtained from the WT and TG mice at baseline, two weeks after induction of STZ-induced DM, five days after administration of anagliptin (before induction of MI), and four weeks after induction of MI ($n = 5-8$). As shown in Figure 2, DPP4 enzyme activity was increased in the diabetic state and was decreased by administration of anagliptin in both the WT and TG mice, in a similar pattern. These results showed that a dose of 300 mg/kg/day of anagliptin adequately inhibited DPP4 activity in both the WT and TG mice.

Measurements of plasma HMGB1 level: We next examined the plasma levels of HMGB1 using the ELISA kit in order to verify our hypothesis that HMGB1 is one of the

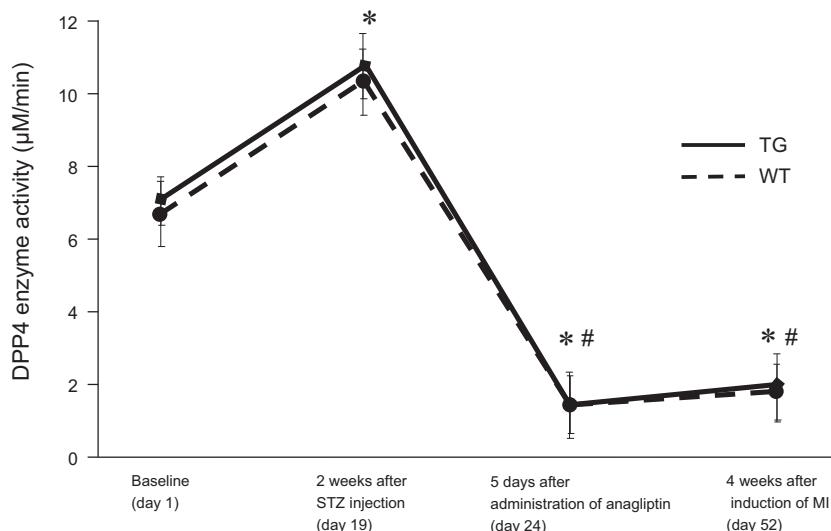


Figure 2. Plasma DPP4 enzyme activity of WT and HMGB-1-TG mice. Representative evaluations of DPP4 enzyme activity at baseline, two weeks after induction of STZ-induced DM, five days after administration of anagliptin, and four weeks after induction of MI in WT and TG mice. The number of mice in each group was 5-8. * $P < 0.05$ versus baseline, # $P < 0.05$ versus two weeks after STZ injection in the same group.

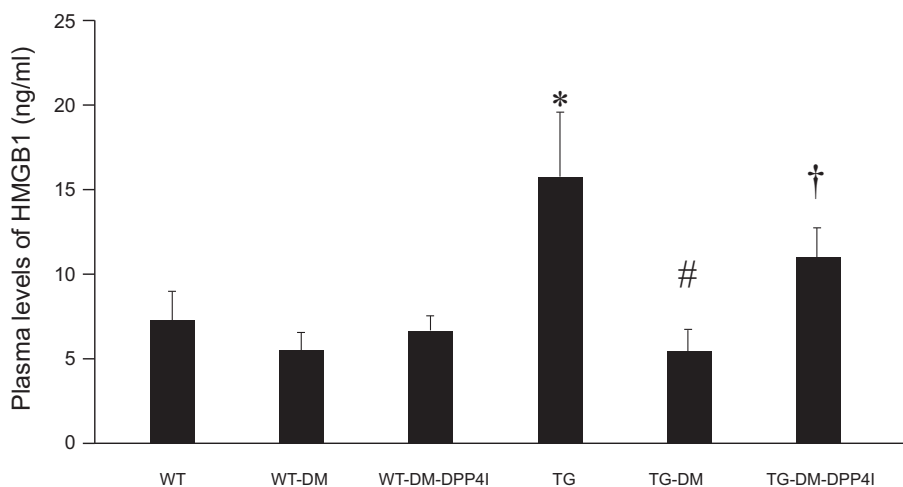


Figure 3. Plasma levels of HMGB1 at four days after induction of MI. The number of mice in each group was 8-10. * $P < 0.05$ versus WT group, # $P < 0.05$ versus TG group, † $P < 0.05$ versus TG-DM group.

potential substrates degraded by DPP4 enzyme, and that inhibition of DPP4 activity increases the plasma levels of HMGB1, resulting in cardio-protective effects in the diabetic state (Figure 3). At four days after induction of MI, HMGB1 plasma levels were significantly higher in the TG group than in the WT group (15.7 ± 3.8 ng/mL versus 8.4 ± 1.6 ng/mL, $P < 0.05$). The TG-DM group showed lower plasma HMGB1 levels (5.5 ± 1.4 ng/mL) than the TG group ($P < 0.05$), and this decline was restored by inhibition of DPP4 activity (10.1 ± 2.2 ng/mL) ($P < 0.05$). No significant differences were observed in the plasma levels of HMGB1 among the WT, WT-DM, and WT-DM-DPP4I groups.

Gravimetric and echocardiographic data at four weeks after MI induction: The gravimetric and echocar-

diographic data at four weeks after induction of MI among the six groups are shown in the Table. Blood sugar (BS) was significantly higher in the DM and DM-DPP4I groups than in the non-DM groups ($P < 0.01$) in both the WT and TG mice. We used a type 1 DM model induced by STZ in the present study, so BS was not significantly decreased by administration of the DPP4 inhibitor. Body weight tended to be decreased by the induction of diabetes, and did not significantly change with administration of the DPP4 inhibitor. The ratio of heart weight to body weight did not show any significant differences among the six groups. The echocardiographic data showed that FS was significantly higher in the TG group than in the WT group ($15.5 \pm 0.84\%$ versus $11.3 \pm 0.91\%$) and deteriorated with the diabetic state ($11.9 \pm 0.64\%$, $P < 0.05$);

Table. Comparison of Baseline Characteristics and Echocardiographic Data in Each Group at Four Weeks After MI

Parameter	WT	WT-DM	WT-DM-DPP4I	TG	TG-DM	TG-DM-DPP4I
Baseline characteristics						
BW (g)	27.3 ± 0.71	25.2 ± 1.59	27.1 ± 0.85	27.1 ± 0.76	25.3 ± 1.46	26.7 ± 0.81
BS (mg/dL)	150.2 ± 8.6	420 ± 17.7**	402.6 ± 24.9**	143.9 ± 9.53	418.6 ± 27.8##	389.7 ± 24.3##
HW/BW (mg/g)	8.08 ± 0.24	7.99 ± 0.23	7.94 ± 0.34	7.20 ± 0.18	8.29 ± 0.36	7.35 ± 0.33
Echocardiography						
IVST (mm)	0.30 ± 0.04	0.32 ± 0.02	0.33 ± 0.02	0.34 ± 0.04	0.33 ± 0.02	0.33 ± 0.03
PWT (mm)	0.82 ± 0.04	0.76 ± 0.03	0.78 ± 0.05	0.78 ± 0.05	0.85 ± 0.05	0.83 ± 0.05
LVDd (mm)	5.87 ± 0.02	5.88 ± 0.07	5.85 ± 0.08	5.66 ± 0.15	5.90 ± 0.09	5.66 ± 0.04
LVDs (mm)	5.2 ± 0.17	5.1 ± 0.23	5.1 ± 0.15	4.90 ± 0.18	5.00 ± 0.19	4.78 ± 0.07
FS (%)	11.3 ± 0.91	10.5 ± 0.5	11.1 ± 1.24	15.5 ± 0.84*	11.9 ± 0.64#	14.2 ± 0.44†

BW indicates body weight; BS, blood sugar; HW, heart weight; IVST, interventricular septal thickness; PWT, posterior wall thickness; LVDd, left ventricular end-diastolic dimension; LVDs, left ventricular end-systolic dimension; and FS, fractional shortening. Data are presented as mean ± SE. The number of mice in each group was 8-10. * $P < 0.05$ versus WT group, ** $P < 0.01$ versus WT group, # $P < 0.05$ versus TG group, ## $P < 0.01$ versus TG group, † $P < 0.05$ versus TG-DM group.

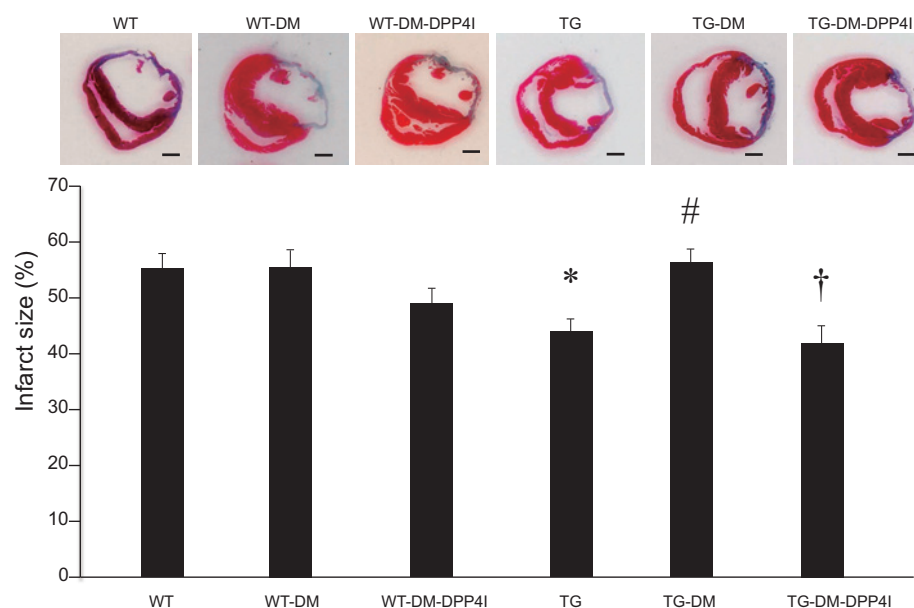


Figure 4. The size of the infarct area at four weeks after induction of MI. Representative photomicrographs show myocardial cross sections stained with Masson's trichrome. The number of mice in each group was 8-10. The scale bar shows 2 mm. * $P < 0.05$ versus WT group, # $P < 0.05$ versus TG group, † $P < 0.05$ versus TG-DM group.

however, it was ameliorated by the inhibition of DPP4 activity ($14.2 \pm 0.44\%$, $P < 0.05$).

Assessment of infarct size: We then investigated the impact of DPP4 inhibition on the infarct area at four weeks after induction of MI (Figure 4). The size of the infarct area was significantly smaller in the TG mice than in the WT mice ($44.8 \pm 2.4\%$ versus $55.2 \pm 2.6\%$, $P < 0.05$) as previously reported.¹²⁾ The TG-DM group had a significantly broader infarct area compared with the TG group ($56.3 \pm 2.4\%$) ($P < 0.05$). The size of the infarct area was reduced by the inhibition of DPP4 enzyme activity ($41.8 \pm 3.4\%$) ($P < 0.05$). No significant differences were observed among the WT, WT-DM, and WT-DM-DPP4I groups.

Evaluation of capillary and arteriole density in the MI border zone: We evaluated the number of PECAM-1-

and α SMA-positive cells in the MI border zone as indicators of angiogenesis (Figures 5, 6). As shown in Figure 5, capillary density, defined by the number of PECAM-1-positive cells in the MI border zone, was higher in the TG group than in the WT group (54.4 ± 2.3 versus $38.2 \pm 2.1/\text{HPF}$, $P < 0.05$). In the TG mice, the capillary density in the MI border zone was significantly reduced by induction of DM ($40.9 \pm 3.2/\text{HPF}$, $P < 0.05$); however, this reduction was restored to levels nearly equal to those of the TG group by administration of the DPP4 inhibitor ($53.1 \pm 3.8/\text{HPF}$). There were no significant differences in capillary density among the WT, WT-DM, and WT-DM-DPP4I groups (Figure 5).

Arteriole density was assessed by the number of α SMA-positive cells, which were smooth muscle cells of the arterioles in the MI border zone. Arteriole density had

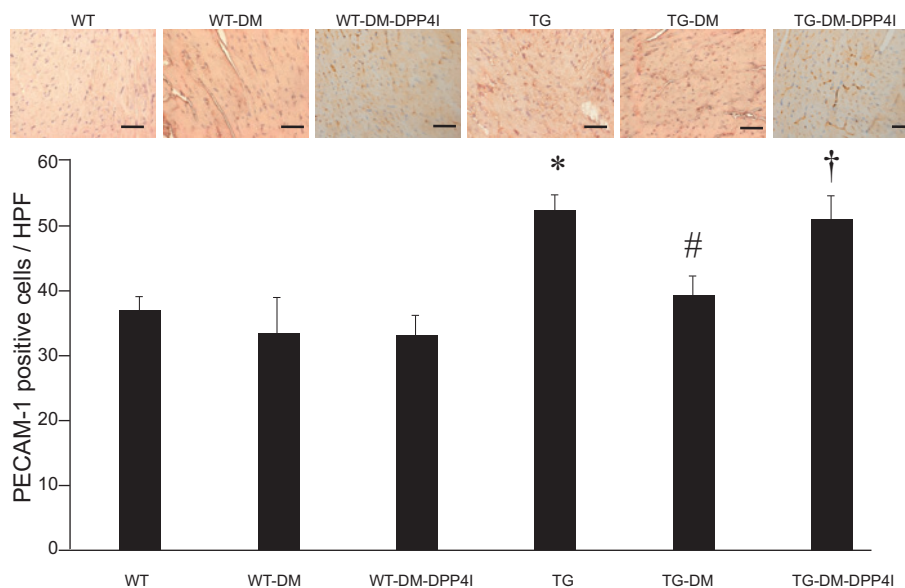


Figure 5. Capillary density in the border zone at four weeks after induction of MI. Representative photomicrographs show myocardial tissue with immunohistochemical staining with anti-PECAM-1 antibody. PECAM-1-positive cells indicate microvascular endothelium. The number of mice in each group was 7-9. The scale bar shows 50 μ m. * $P < 0.05$ versus WT group, # $P < 0.05$ versus TG group, † $P < 0.05$ versus TG-DM group.

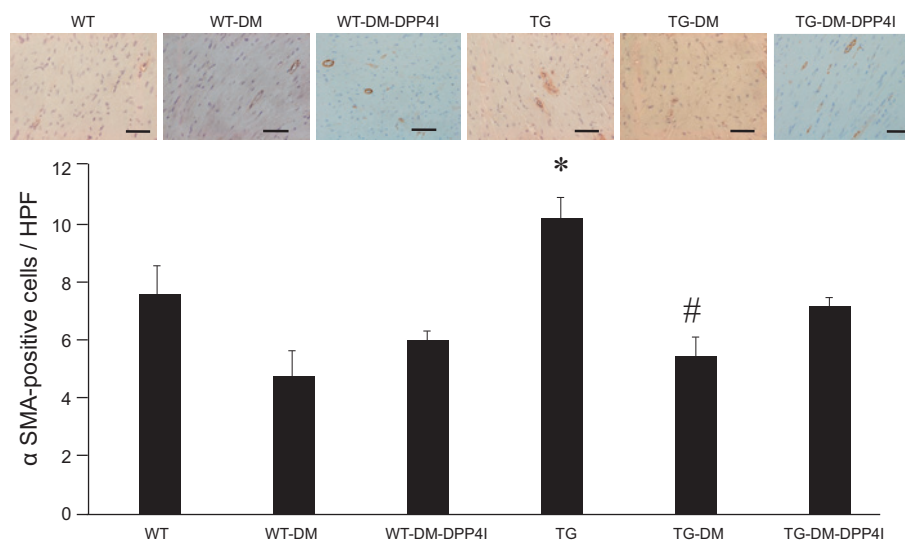


Figure 6. Arteriole density in the border zone at four weeks after induction of MI. Representative photomicrograph shows myocardial tissue with immunohistochemical staining with anti- α SMA antibody. The number of mice in each group was 6-9. The scale bar shows 50 μ m. * $P < 0.05$ versus WT group, # $P < 0.05$ versus TG group.

similar tendencies to capillary density, as both were reduced by DM and restored by DPP4 inhibition. Briefly, arteriole density was reduced by DM and restored by the DPP4 inhibition (Figure 6).

Evaluation of signaling promoting angiogenesis: HMGB1 is known to promote tissue angiogenesis via VEGF.^{14,21)} To investigate the mechanism of such promotion by DPP4 inhibition in diabetic TG mice, we analyzed the expression of VEGF in the MI border zone by West-

ern blotting (Figure 7). VEGF expression was promoted by the induction of MI in both the WT and TG mice in the MI border zone compared with sham-operated mice, although this increase was not statistically significant in the WT mice. The TG mice had significantly higher VEGF expression in the MI border zone than the WT mice ($P < 0.05$). VEGF expression in the MI border zone was significantly suppressed in the TG mice with diabetes and was ameliorated by inhibition of DPP4 activity ($P <$

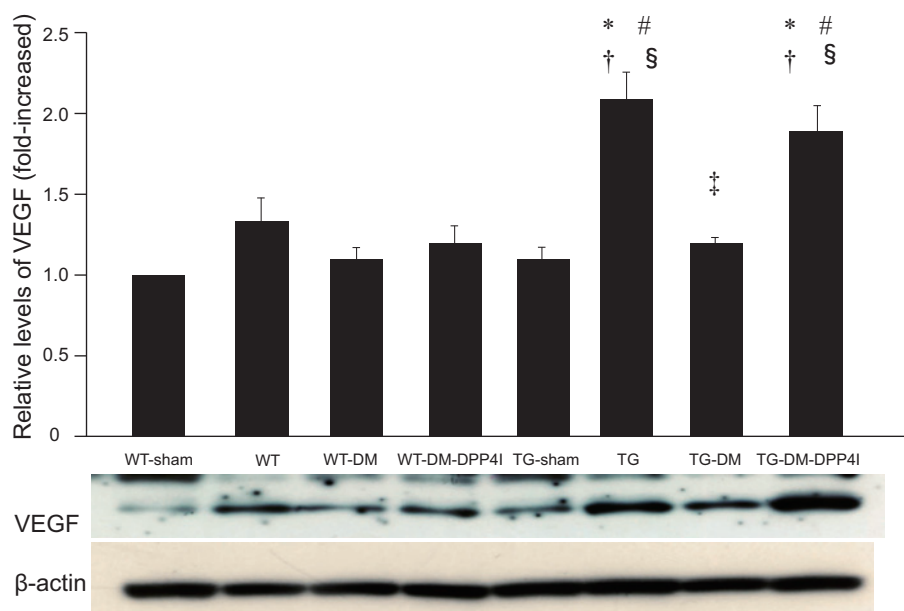


Figure 7. VEGF expression in MI border zone. Representative figure shows Western blotting of VEGF protein in MI border zone at four days after induction of MI. The number of mice in each group was 6. Data are represented as relative values to the WT-sham group. * $P < 0.05$ versus WT-sham, # $P < 0.05$ versus WT, † $P < 0.05$ versus TG-sham, § $P < 0.05$ versus TG, § $P < 0.05$ versus TG-DM.

0.05).

Discussion

This study demonstrated that the function of HMGB1 in promoting angiogenesis was reversed in the diabetic state. We additionally showed that DPP4 inhibition suppressed the degradation of HMGB1, resulting in the promotion of tissue angiogenesis and amelioration of cardiac function after MI. These findings indicate that HMGB1 was possibly one of the substrates of DPP4, and that the inhibition of DPP4 activity led to the preservation of cardiac function after MI via HMGB1.

HMGB1 is a non-histone DNA-binding protein consisting of 215 amino acids, and ubiquitously exists in every cell type.^{5-7,28} Intracellular HMGB1 plays important roles in regulating genome replication, transcription, and DNA repair.^{29,30} HMGB1 also has a crucial role in survival because HMGB1-gene knockout mice die within the first day of life due to hypoglycemia.³¹ HMGB1 released from necrotic cells (extracellular HMGB1) acts as a cytokine involved in inflammation through the receptor for advanced glycation end products and toll-like receptors (TLR) 2 and 4.^{32,33} One of the local actions of extracellular HMGB1 is stimulation of tissue repair by recruiting stem cells and promotion of their proliferation.¹¹ With regard to the function of HMGB1 in tissue regeneration, Kitahara, *et al.* reported that HMGB1 restores cardiac function through the activation of tissue angiogenesis after MI using male mice with a four-fold expression of HMGB1 in cardiac muscle cells (TG mice).¹² Based on Kitahara, *et al.*'s report, we demonstrated that HMGB1-induced angiogenesis after MI arose from homing and differentiating endothelial progenitor cells from bone marrow, resulting

in promoting angiogenesis at the peri-infarct area.¹⁴ These reports showed that HMGB1 was released from necrotic myocardial cells into circulation, and plasma HMGB1 level after MI was much higher in TG mice than in WT mice. With regards to angiogenesis, HMGB1 promotes tissue angiogenesis by increasing the production of pro-angiogenic cytokines including VEGF, TNF- α , and IL-8 from endothelial cells and macrophages.^{34,35}

DPP4, a type of peptidase and one of the therapeutic targets for DM, reportedly degrades and inactivates not only incretin hormones but also other substances, including SDF-1, BNP, and substance P.¹⁹ Because HMGB1 has a component of partial DPP4 cleavage sites,²⁰ it can also be degraded and inactivated by DPP4 *in vivo*. Straino, *et al.* reported that HMGB1 levels of mice skin tissue were decreased in the diabetic state, and HMGB1 application to skin wounds promoted the tissue angiogenesis and wound healing.³⁶ Furthermore, Marchetti, *et al.* proved that HMGB-1 was degraded by DPP4 in *in vitro* experiments and the inhibition of DPP4 enhanced tissue angiogenesis in a murine skin-wound model.²⁰

In the present study, we firstly demonstrated *in vivo* that cardio-protective effects, such as promotion of angiogenesis, suppression of cardiac remodeling, and enhancement of VEGF expression in the peri-infarct area, were reversed in diabetic state and ameliorated by administration of DPP4 inhibitor in the TG mice. However, these reactions could not be observed in the WT mice. In WT-DM mice, slight decrease of plasma HMGB1 level was observed compared to non-DM WT mice, although this was not statistically significant (Figure 3). Additionally, in Figure 6, the number of α SMA positive cells was lower in WT-DM mice than in non-DM WT mice, and that was partially recovered by the DPP4 inhibition, although these

differences were not statistically significant. Taken together, similar trends with TG mice were observed in non-DM WT, WT-DM, and WT-DM-DPP4 inhibition groups, but these levels were low, and thus did not lead to decrease in infarct size in WT mice. Sauve *et al.* reported that infarct size in genetic DPP4 deletion mice did not differ with that in WT mice,³⁷⁾ and their data were consistent with our present study. In addition, Shigeta, *et al.* reported that VEGF expression in cardiac muscle tissue was suppressed in the diabetic state with inhibition of DPP4 activity in a chronic heart failure model induced by trans-aortic constriction.³⁸⁾ In the current study, VEGF expression was escalated in diabetic HMGB1-TG mice by DPP4 inhibition compared with that in diabetic HMGB-1 TG mice without inhibition of DPP4. This indicates that HMGB1 is a substrate of DPP4 *in vivo*, and is inactivated under diabetic conditions in which DPP4 activity increases, resulting in deterioration of cardiac function after MI in HMGB1-TG mice.

Although the DPP4 inhibition showed favorable effects on heart failure in several basic researches, recent major clinical trials (SAVOR, EXAMINE, TECOS) suggested that DPP4 inhibitors were not associated with decreases of major adverse cardiac events in DM patients with a history of established cardiovascular diseases or multiple cardiovascular risk factors.³⁹⁻⁴¹⁾ In addition, the secondary analysis of SAVOR trial indicated DPP4 inhibitor, saxagliptin, increased the hospitalization of worsening heart failure.⁴²⁾ However, these results did not deny the cardio-protective effects of DPP4 inhibition from some reasons. First, these trials were placebo-controlled trials carried out to confirm the safety of DPP4 inhibitors on patients with cardiovascular diseases, not designed to demonstrate beneficial effect of DPP4 inhibitor on heart failure. Moreover, it is also possible that observational periods of these trials were too short to reveal the cardiac benefit of DPP4 inhibitors. Second, DPP4 inhibitors might have class effects on heart failure. Whereas subgroup analysis of SAVOR trial showed unfavorable cardiac effect of DPP4 inhibitor,⁴²⁾ EXAMINE and TECOS showed non-inferiority of DPP4 inhibitors on hospitalization of heart failure.^{43,44)} In addition, subgroup analysis of EXAMINE trial showed DPP4 inhibitors decreased cardiovascular mortality compared with placebo in the highest BNP quartiles.⁴³⁾ Therefore, the impact of DPP4 inhibitor on cardiovascular diseases has remained unclear, and further clinical studies should be conducted to clarify the effect of DPP4 inhibitors on cardiovascular diseases and heart failure.

This study includes several limitations. First, since DPP4-inhibitors are generally used for treatment of type 2 DM, further studies are needed to verify cardio-protective effect of the DPP4 inhibition via HMGB1 in type 2 DM model mice. Second, we demonstrated cardio-protective effect of the DPP4 inhibition in only diabetic model in present study. Further study to examine the effect of the DPP4 inhibition in non-DM model is required. Third, the induction of experimental myocardial infarction might have influenced on DPP4 activity. However, our essential purpose of this study was to examine whether the DPP4 inhibition restores HMGB1-induced angiogenesis and tis-

sue repair in diabetic mice after MI. A recent article by Kubota and Takano, *et al.*⁴⁵⁾ has demonstrated that the treatment with DPP4 inhibitor increased the ratio of endothelial cell numbers to a cardiomyocyte, improved cardiac function and decreased the infarct size in C57BL/6 mice after MI, suggesting a key role of DPP4 activity after MI in non-diabetic condition. Thus, we should consider effects of DPP4 activity on HMGB1 function in non-diabetic condition in the future study.

Conclusion

We demonstrated that the inhibition of DPP4 activity blocked HMGB-1 degradation, resulting in angiogenesis promotion and amelioration of cardiac function after myocardial infarction in diabetes.

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Disclosures

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